

Review

The p38 MAP Kinase Family as Regulators of Proinflammatory Cytokine Production in Degenerative Diseases of the CNS

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ABSTRACT: Inflammation in the central nervous system (CNS) is a common feature of age-related neurodegenerative diseases. Proinflammatory cytokines, such as IL-1 β and TNF α , are produced primarily by cells of the innate immune system, namely microglia in the CNS, and are believed to contribute to the neuronal damage seen in the disease. The p38 mitogen-activated protein kinase (MAPK) is one of the kinase pathways that regulate the production of IL-1 β and TNF α . Importantly, small molecule inhibitors of the p38 MAPK family have been developed and show efficacy in blocking the production of IL-1 β and TNF α . The p38 family consists of at least four isoforms (p38 α , β , γ , δ) encoded by separate genes. Recent studies have begun to demonstrate unique functions of the different isoforms, with p38 α being implicated as the key isoform involved in CNS inflammation. Interestingly, there is also emerging evidence that two downstream substrates of p38 may have opposing roles, with MK2 being pro-inflammatory and MSK1/2 being anti-inflammatory. This review discusses the properties, function and regulation of the p38 MAPK family as it relates to cytokine production in the CNS.

Key words: Protein kinase; Neuroinflammation; Neurodegeneration; Microglia; Signal Transduction

One of the primary classes of inflammatory mediators throughout the body, including the central nervous system (CNS), is cytokines. Cytokines are signaling molecules that act through specific receptors and signal transduction pathways to exert a particular biological response in a target cell. Extensive evidence from both clinical studies and preclinical animal models has implicated overproduction of proinflammatory cytokines as a contributor to pathophysiology progression in chronic neurodegenerative disorders like Alzheimer's disease (AD), Parkinson's disease (PD), and multiple sclerosis (for review, see [1]). Several studies have also documented a key role for proinflammatory cytokine overproduction as a potential driving force for pathology progression in acute neurodegenerative disorders such as traumatic brain injury and stroke.

Taken in its entirety, the evidence is consistent with the hypothesis that proinflammatory cytokine up-regulation is a comparatively early event in the progression of pathophysiology that is causally linked to synaptic dysfunction, behavior deficits and, in the more extreme case, neuronal death. These data also suggest that strategies to target signaling pathways that lead to cytokine overproduction should be explored in attempts to develop new CNS therapeutics with the potential for disease modification in multiple diseases and clinical presentations.

A key signal transduction pathway involved in the production of proinflammatory cytokines is the p38 mitogen-activated protein kinase (MAPK) pathway (Figure 1). Interestingly, p38 MAPK is also one of the primary signal transduction pathways activated by the cytokines IL-1 β and TNF α [2-5]. This raises the

potential for a specific p38 MAPK-activating stimulus to lead to a state where inflammation begets inflammation. In the AD brain, for example, the initial stimulus could be an increase in the protein amyloid-beta ($A\beta$), which can activate p38 MAPK and proinflammatory cytokine release from microglia. Excessive proinflammatory cytokine production can then lead to further p38 MAPK activation in microglia or other CNS cell types (e.g., astrocytes and neurons) and a resultant potential feedback loop that exacerbates disease. The p38 MAPK pathway, therefore, appears to be a critical nexus between pathological microglial activation and detrimental inflammation in CNS disorders. If this link can be disrupted selectively, balance may be able to be restored.

The goal of this review is to present an overview of the p38 MAPK family as it relates to CNS proinflammatory cytokine production. We provide a basic introduction to the p38 MAPK family that emphasizes the role of individual p38 MAPK isoforms in the production of proinflammatory cytokines. We also discuss various approaches used to implicate particular p38 MAPK isoforms, especially p38 α , in cytokine production. Finally, we discuss the interface of p38 MAPK with two of its downstream substrates in the regulation of proinflammatory responses. The overall purpose is to explore the evidence that p38 MAPK signaling pathways may be viable targets for modulating inflammatory responses in neurological diseases.

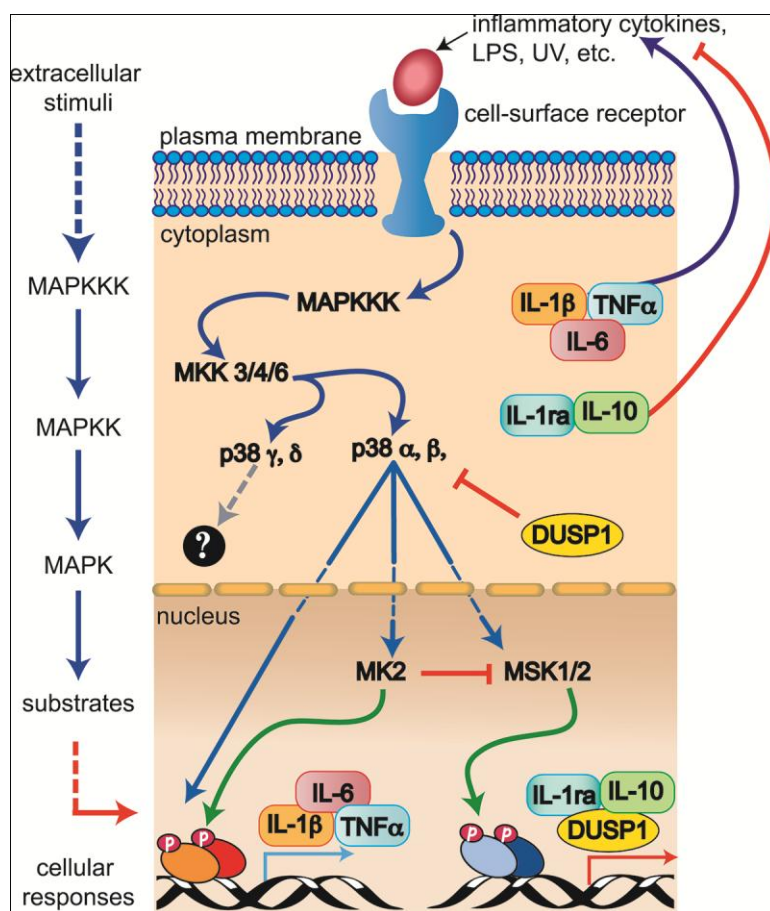


Figure 1: Overview of the p38 MAPK signal transduction pathway involved in cytokine production. Extracellular stimuli activate receptor and receptor associated proteins, which then transduce the activating signal into cellular responses by a three step MAPK pathway (MAPKKK, MAPKK, MAPK) that leads to p38 activation. Activated p38 can directly activate transcription factors or can act indirectly via two downstream kinases MK2 and MSK1/2. The cellular response produced can have many paracrine functions but can also have autocrine functions leading to further activation or suppression of the p38 pathway.

The p38 family of protein kinases

The serine-threonine protein kinase p38 was identified as a 38kDa protein that is phosphorylated within 15

minutes in response to lipopolysaccharide (LPS) [5, 6]. Subsequent studies demonstrated that p38 (later named p38 α) was strongly activated by UV radiation,

and also by the proinflammatory cytokines, IL-1 β and TNF α [2-4]. The p38 MAPK family consists of four major isoforms encoded by separate genes: p38 α (MAPK14), p38 β (MAPK11), p38 γ (MAPK12), and p38 δ (MAPK13). These isoforms are expressed in different tissues and have distinct functions. In a general sense, the p38 MAPKs integrate and process extracellular stimuli through a series of intracellular signaling complexes and phosphorylation cascades that lead to stimulus-specific responses. In response to inflammatory stimuli, p38 MAPK is activated by a series of upstream kinases that lead to dual phosphorylation of Thr and Tyr residues in p38. The activated p38 MAPK then phosphorylates appropriate downstream substrates that culminate in increased production of proinflammatory cytokines by a number of different mechanisms (reviewed in [7]).

p38 α MAPK and p38 β MAPK

The α and β isoforms of p38 MAPK share 75% homology [8]. The expression of p38 α and p38 β is highest in the heart, skeletal muscle and brain, but there are low levels of expression of these two isoforms in most tissues [8, 9]. Most stimuli that activate p38 α also activate p38 β to a similar degree, although exceptions have been reported. For example, in HeLa cells transfected with either p38 α or p38 β [8], TNF α and IL-1 β both activate p38 α . However, TNF α stimulates p38 β activity to a much larger degree than does IL-1 β . Furthermore, the response kinetics to TNF α are different; p38 α is quickly activated within 15 minutes after stimulation, whereas the response by p38 β is slower and slightly blunted [8]. The most striking difference was a strong preference by p38 β for one of its substrates, ATF2. Overexpression of p38 β increased ATF2 reporter gene expression ~20-fold better than p38 α [20]. In addition, p38 β has ~180-fold higher catalytic activity for ATF2 than does p38 α [8,10]. Although the differences between p38 α and p38 β may be subtle, it is clear that these isoforms may be regulated differently and work via different downstream substrates in response to specific inflammatory stimuli.

Insight into the role of p38 α MAPK and p38 β MAPK in cytokine responses

A number of different approaches have been used to explore the role of p38 MAPK in inflammatory cytokine production. The pyridinyl imidazole class of compounds was one of the first small molecule

inhibitors of p38 MAPK, and have been excellent tools in validating the importance of p38 MAPK in inflammatory cytokine production. However, these early generation p38 MAPK inhibitors were not very selective for p38 and also did not distinguish between p38 α and p38 β [8], limiting the ability to infer specific functions of the individual p38 isoforms. To further hamper the understanding of the function of the p38 α isoform, global deletion of the p38 α gene in mice is embryonically lethal, due to a requirement for p38 α in placental development and organogenesis [11-14]. On the other hand, mice null for the p38 β gene have no known developmental phenotype [15]. Despite these early impediments, newer approaches using drug-resistant knock-in mice, tissue-selective conditional p38 α gene deletions, and new classes of more selective inhibitors have provided increased insight into distinct functions of p38 α and p38 β .

Drug-resistant, knock-in mice

The gatekeeper residue of protein kinases is a key residue in the ATP binding fold whose side chain is situated in front of a conserved hydrophobic pocket region, but is not required for kinase activity. In most protein kinases, the gatekeeper residue is an amino acid with a bulky side chain such as Met, Leu, or Phe that will block access to this pocket [16]. However, in p38 α and p38 β , the gatekeeper residue is an amino acid with a small side chain (Thr-106), which allows entrance to this hydrophobic region. Many p38 inhibitors utilize the accessibility of the hydrophobic pocket for specificity for the p38 α and p38 β isoforms over the p38 γ /p38 δ isoforms which have a larger Met gatekeeper residue. In addition, many inhibitors including the pyridinyl imidazole based inhibitors [17, 18] utilize stabilizing hydrogen bond interactions with the amide nitrogen of Met-109 for increased inhibitor affinity.

O'Keefe et al [17, 18] exploited the kinase gatekeeper selectivity filter to generate a point mutant of either p38 α or p38 β where Met is substituted for Thr at position 106 (T106M). The mutant kinase retains full kinase activity, but is no longer susceptible to inhibition by small molecule inhibitors that utilize this structural feature. They then generated mouse strains that have either a knocked-in p38 α (T106M) or p38 β (T106M) whose insertion disrupts the endogenous p38 isoform. These mice retain full p38 kinase activity, but the enzyme can no longer bind to the p38 inhibitors. For the first time the question could be asked *in vivo* which p38 isoform, α or β , was

responsible for the effects of p38 inhibitors. The answer: p38 α .

In the p38 α (T106M) mice, O'Keefe and colleagues [17,18] demonstrated that it was indeed the p38 α form of the kinase that was important for cytokine production. Treatment of either wild type mice or p38 β (T106M) mice with a p38 inhibitor was equally effective at blocking LPS-induced increases in TNF α in the plasma. However, in the drug resistant, p38 α (T106M) mice, the p38 inhibitor was completely ineffective [18]. This nicely designed study, for the first time, clearly showed that selective inhibition of the p38 α isoform *in vivo* is sufficient and necessary for suppression of peripherally increased proinflammatory cytokine levels after an inflammatory challenge. Whether similar results occur in CNS inflammatory conditions has not been reported yet, but these knock-in mice will provide valuable tools for validating that new CNS-penetrant small molecule cytokine suppressors are exerting their CNS effects via inhibition of p38 α .

Conditional p38 α knock-out mice

The second approach to study of p38 α is the use of tissue-selective, conditional knock-out (KO) mice. This approach has been very fruitful in enhancing our understanding of what p38 α does in different cell types. In 2008, Kang and colleagues [19] developed a mouse that was null for p38 α only in cells of the monocyte/macrophage lineage (p38 α ΔM). Following either LPS stimulation or cecal ligation and puncture, p38 α ΔM mice were protected from the lethal effects of both LPS and cecal ligation and puncture induced sepsis. The protective effect appeared to be due to a reduced TNF α response, which in both sepsis models was almost absent in the p38 α ΔM mice [19]. Another *in vivo* model that has been tested in the p38 α ΔM mice is a skin inflammation model. This tissue injury model produced some unexpected findings. Despite the protection that had been shown against septic shock, the p38 α ΔM mice had a worse outcome in the skin inflammation model. There are many possible explanations for why in this model the injury was exacerbated, but two responses that may have contributed were an increase in COX-2 expression and a decrease in the anti-inflammatory cytokine IL-10 in the p38 α ΔM mice compared to wild type mice [20].

Kim et al. [20] took the study of the p38 α ΔM mice a step further, to implicate another potential mechanism for the increased inflammation. There are multiple points in the p38 MAPK signaling pathways

where regulation can occur. Points of negative regulation include receptor desensitization, dissociation or deactivation of signaling pathway mediators, and kinase dephosphorylation by phosphatases. Dephosphorylation of some protein kinases is a fast and efficient way of shutting down kinase activity [21]. One phosphatase that is important for dephosphorylation of p38 MAPK is dual-specificity protein phosphatase (DUSP)1 (also known as MKP-1). DUSP1 is an immediate-early gene which is involved in the dephosphorylation of p38 and JNK, and also to a lesser extent ERK [21]. Macrophages isolated from DUSP1 deficient mice have sustained activation of p38, and increased TNF- α following LPS [22]. The temporal activation of DUSP1 occurs shortly after that of p38. For example, following LPS stimulation, DUSP1 protein becomes elevated at 45 minutes and peaks at 60 minutes. In contrast, phospho-p38 expression peaks at 15 minutes and reaches baseline at 45 minutes [22]. In DUSP1 deficient macrophages, LPS-induced activation and subsequent deactivation of phospho-p38 were significantly slowed, resulting in prolonged duration of action [23]. Importantly, a dose of LPS that was normally sublethal in wild type mice was 80% lethal in DUSP1 deficient mice by 72 hours [23], demonstrating that DUSP1 plays a role in limiting p38 activation.

An interesting observation is that p38 α activation and signaling can also drive DUSP1 production, thus providing a mechanism for negative feedback to limit p38 activation [20]. The p38 α null macrophages showed a significantly diminished induction of DUSP1 in response to LPS. In addition, as DUSP1 also dephosphorylates JNK and ERK, the decrease in DUSP1 seen in the p38 α ΔM resulted in an increase in phospho-JNK and phospho-ERK. Thus, the increased inflammation seen in the p38 α ΔM may result from both inadequate DUSP1 production as well as increases in other MAPK pathways (JNK and ERK) that may converge on inflammatory responses.

p38 α -selective small molecule inhibitors

The third approach to study the function of p38 α is through the use of newer, more isoform selective p38 inhibitors that have been designed using the extensive crystallographic data available on p38 structures. This is also the only approach of the three that has been used to explore the role of p38 α in brain injury and neurodegenerative disease, and there are two examples of the effectiveness of the p38 α specific

inhibitors in preventing neurodegeneration. The first example was shown using the second-generation p38 α inhibitor, SB239063, in the middle cerebral artery occlusion (MCAO) model of brain injury. There was a reduction in the infarct volume following MCAO in rats that received the small molecule p38 α inhibitor, as well as a lessening of the neurological deficits following injury [24, 25].

A second example implicating p38 α in CNS disorders was demonstrated in an AD-relevant mouse model. A selective, CNS-penetrant p38 α inhibitor, MW01-2-069A-SRM, was developed and used by Munoz et al. [26] in an amyloid-beta (A β)-induced model of brain injury. MW01-2069A-SRM was found to block the A β -induced increase in levels of the proinflammatory cytokines TNF α and IL-1 β in the hippocampus. The p38 α inhibitor also prevented synaptic dysfunction and hippocampal-dependent behavioral deficits caused by the A β insult [26]. These studies indicate that p38 α could be a credible target for drug development campaigns for AD and related neurodegenerative diseases where overproduction of cytokines is involved in the disease mechanisms.

A recent study highlighted a potential caveat to the use of non-selective p38 α /p38 β inhibitors. Greenblatt et al. [27] found that in p38 β null mice, which previously have been reported to have no phenotype, there were significant skeletal abnormalities including a thinning of the bones. There was also some thinning in the p38 α conditional KO mouse, but the effect was less pronounced. The thinning of the bones was also seen after treatment with the non-selective p38 α /p38 β inhibitor SB203580 [27]. The authors suggest that p38 β agonists may help prevent osteoporosis. Development of selective p38 α inhibitors that spare p38 β might need to be considered for an aging population where osteoporosis can be a complication.

p38 γ MAPK and p38 δ MAPK

The highest expression of p38 δ is limited to the lungs and kidneys [28]; yet, in a very intriguing study, Hale et al. [29] investigated the expression of the four p38 MAPK isoforms in different immune cells. The p38 α isoform was widely expressed, and was the dominant p38 isoform in the immune cell study. Endothelial cells were the one exception. In endothelial cells, lower levels of p38 α were found and there was robust expression of p38 β . The interesting finding concerning p38 δ resulted from a comparison of monocytes and macrophages. In monocytes, the

precursor to dendritic cells and macrophages, there were high levels of p38 α and only low levels of p38 δ . However, in monocytes that had been differentiated into macrophages *in vitro*, the levels of p38 α were reduced compared to the levels in monocytes. As determined by real-time PCR, in this population of macrophages, p38 δ became the dominant p38 isoform, with ~4-fold increased expression compared to p38 α [29].

Fearn et al. [30] demonstrated using monocyte cell lines and an astrocytoma cell line that IL-1 β and TNF α , or TLR-2 or TLR-4 ligands could activate all four p38 family members. The degree of activation of the different p38 family members was cell and stimulus specific. For example, in the astrocytoma cell line, IL-1 β caused the greatest induction in p38 β , p38 γ , and p38 δ , whereas LPS caused the greatest increase in p38 α . In the THP1- wtCD14 monocyte cell line, the TLR-2 ligand caused a large induction of all 4 isoforms, whereas the cytokines activated p38 α , but not p38 β , p38 γ , or p38 δ [30]. The expression of p38 δ has not been looked at in microglia. It is also not known if there are changes in the expression of p38 δ in neurodegenerative disease.

The p38 δ (MAPK13) isoform shares 60% homology with the other p38 family members [28], and can be acted on by the upstream MAPKKK and MAPKK (MKK3, MKK4, and MKK6) kinases that activate other p38 family members [28]. This suggests that p38 δ should be activated by stimuli that activate the other p38 family members. This was confirmed by demonstrating activation of p38 δ kinase activity after stimulation by anisomycin, UV radiation and TNF α that was similar to that of p38 α kinase activation. However, IL-1 β and IL-6 were found to be weaker activators of p38 δ compared to p38 α [28]. In response to LPS, macrophages exhibited an increase in a 41kDa phospho-p38 band which represented phospho-p38 α , and a 43.5kDa band which represented phospho-p38 δ [29]. This separation into phospho-p38 α and phospho-p38 δ could also be seen in PC12 cells following hypoxia [31]. In COS-7 cells transfected with tau and each of the four p38 isoforms, it was found that only p38 γ and p38 δ could efficiently phosphorylate tau [32]. As p38 γ and p38 δ are not the major p38 isoform in neurons, it is unclear how relevant these results are to Alzheimer's disease. In a subsequent study [33], it was found that all four p38 isoforms could phosphorylate tau, that Thr50 in tau was an important site of phosphorylation by p38, and that phosphorylation of tau at this residue was most

dependent on p38 δ . Finally, the authors showed that phosphorylation of Thr50 was found in Alzheimer's disease brains [33].

The final member of the family, p38 γ (MAPK12), shares 60% homology with p38 α and β [9]. The expression of p38 γ is localized to skeletal muscle. Inflammatory p38 substrates, MK2 and ATF2, are not targets of p38 γ [9] and p38 γ is not thought to be important in inflammation.

MK2: Is this downstream target of p38 the next logical step in anti-cytokine therapeutics?

Mitogen-activated protein kinase-activated protein kinase 2 (MK2) is one of several kinases activated through direct phosphorylation by p38 [34]. MK2 kinase activity is dependent on activation by p38 α . In p38 α negative embryonic stem cells (ES) cells treated with strong p38 activators, anisomycin (50 ng/ml) or sodium arsenite (0.5 mM), there was essentially no MK2 kinase activity [12]. The expression of p38 and MK2 are each reliant on the other, as loss of MK2 causes a decrease in the amount of total p38 [35].

In 1999, Kotlyarov and colleagues [36] developed a MK2 KO mouse and established the importance of MK2 in regulating TNF α . Upon high dose LPS stimulation, which was lethal in wild type mice, the MK2 KO mice were found to be resistant to the LPS injury. The reasons for the resistance were that there was only a very limited cytokine storm in response to the LPS, and there was a decreased induction of a number of cytokines including IFN- γ , IL- β IL-6, and IL-10. In addition, there was essentially no up-regulation of TNF α in the serum of these mice, as well as a dramatic decrease in nitric oxide levels [36]. MAP kinase-activated protein kinase 3 (MK3) is 70% identical in structure to MK2. Lack of MK3 alone does not produce an altered cytokine response; however, the absence of both MK2 and MK3 does produce a further decrease in TNF α from the levels of the MK2 KO alone [37]. MK2 was subsequently shown to directly regulate TNF α and IL-6 at the 3'-untranslated region (UTR) of mRNA in the AU-rich elements (AREs) and thereby regulate mRNA stability [38]. Interestingly, MK2 also inhibits the phosphorylation of MSK1/2 by p38 [39] (See figure 1 for simplified overview of interactions). In the CNS, the expression of MK2 can be found in neurons, astrocytes, and microglia; however, like p38 α , the expression of MK2 in microglia is 5 to 10 fold higher than in neurons or astrocytes [40]. In microglia

stimulated with LPS and IFN- γ , 75% of the increase in TNF α can be blocked by MK2 genetic deletion [40]. A reduction in TNF α , IL-6, KC and nitric oxide was found in mixed neuron-glia cultures treated with LPS from MK2 KO mice compared to wild type mice [41].

The effects of MK2 deficiency have been investigated in a number of neurological disease models, including stroke, AD, and PD. MK2 KO mice were protected from stroke in either a permanent or transient MCAO model, and the loss of MK2 almost completely prevented the up-regulation of IL-1 β that was seen in the wild type mice [42]. In neuron-glia co-cultures treated with A β , there was decreased A β -induced neuronal death when the microglia were from MK2 KO mice [40]. In the MPTP mouse model of Parkinson's disease, there was a sparing of the tyrosine hydroxylase positive neurons in the MK2 KO mice [41]. While there was a decrease in TNF α and nitric oxide expression in the MK2 KO mice following MPTP, there was no change in the microglia activation as measured by MAC-1 expression [41]. Decreased TNF α responses in Parkinson's disease models have previously been shown to be neuroprotective [43,44]. Therefore, it is likely that the reduced TNF α response in the MK2 KO mice correlates with the neuroprotective effect in the MPTP model [41].

To explore potential mechanisms by which MK2 mice neurons were protected from the MPTP, Thomas et al. [41] used neuron-glia co-culture. Suspecting that the cell death was mediated via nitric oxide, they tested to see if loss of MK2 in the neurons was directly protecting neurons from nitric oxide-induced cell death. Using a chemical nitric oxide donor, 100 μ M sodium nitroprusside, nitric oxide-induced cell death in enriched neuronal cultures was found to be independent of MK2. Interestingly, pretreatment of the neuronal cultures with a p38 inhibitor did protect the neurons from the nitric oxide. The neuronal protection by p38 inhibition was similar for both the MK2 $^{+/+}$ and MK2 $^{-/-}$ neurons. Thus, the neuronal survival effects of p38 inhibition in response to nitric oxide are a result of an MK2 independent mechanism [41].

Therapeutically it appears that there may be different conditions when targeting only MK2 might be useful, in comparison to times when a p38 inhibitor may be indicated. For example, it is clear that many of the anti-cytokine effects of blocking p38 are a result of inhibition of MK2; therefore if the goal were to

block cytokine up-regulation following a traumatic brain injury, for example, an MK2 inhibitor could be useful. However, the MK2 inhibitor would appear to lack some of the direct neuroprotective effects of a p38 inhibitor. Depending on the clinical indication then, it may be advantageous to use a p38 inhibitor that will have an impact on both neurons and glia, or a more specific microglia cytokine suppressor with an MK2 inhibitor.

A complicating factor that must be kept in mind when targeting intracellular kinases is that phosphorylation signaling pathways have considerable crosstalk. There are interconnected loops that contribute both positive and negative feedback and others that modulate the response. For example, it has been reported [39] that MK2 can modulate the NF κ B response, causing a sustained activation of NF κ B by maintaining NF κ B in the nucleus. In the absence of MK2, NF κ B is quickly shut off; however the initial NF κ B response is stronger. The regulation of NF κ B by p38 occurs by the ability of the p38 and NF κ B to move in and out of the nucleus [39]. Thus, there is a complex and interconnected system of cross-talking pathways potentially regulated by p38 levels.

The first p38 inhibitors, the pyridinyl imidazole class of compounds, were described in 1994 [45]. It wasn't until 2005, however, when the first small molecule inhibitor of MK2, an aminocyanopyridine, was reported [46]. Since 2005, a number of MK2 inhibitors have been developed. The carboline analogs class of MK2 inhibitors, developed at Boehringer-Ingelheim Pharmaceuticals, have been shown by crystal structure to interact with the hinge region of MK2 (M138–D142) and bind between p-loop and amino acids V118, T206, L193, and Y194. Unfortunately, the first group of carboline analogs was reported to lack activity in cell based assays [47]. In 2008, a pyrazinoindolone class of MK2 inhibitors was described [48]. The lead compound had a reported IC₅₀ for MK2 kinase activity of 3nM and a cell based IC₅₀ for TNF α at 300nM, but the bioavailability of the compound was not sufficient for TNF α inhibition *in vivo* [48]. However, a number of new MK2 inhibitors have been shown to have the ability to inhibit TNF α *in vivo*. For example, Velcicky et al. [49] modified the pyrrolo-pyrimidinone scaffold to synthesize a 3aminopyrazole based inhibitor. At a dose of 100 mg/kg given orally, the aminopyrazole compound caused a 68% reduction in serum levels of TNF α [49]. Another MK2 inhibitor, a β -carboline derivative, was synthesized as an ester prodrug [50]

and resulted in an 84% inhibition of TNF α when administered at 40mg/kg IP in a rat LPS model.

So far, the MK2 inhibitors are still in the development phase. There is only limited efficacy of a small subset of the inhibitors using *in vivo* models. As of the writing of this review, there have been no tests of MK2 inhibitors in disease relevant models such as rheumatoid arthritis. There have been no reported studies in CNS disorders or even information available on whether the compounds are brain-penetrant. Therefore, although MK2 would appear to be a good therapeutic target for CNS disorders, more work is needed. In the last five years, much progress has been made in identifying novel small molecules that have the ability to inhibit MK2. Effort must be expended now to refine these compounds to improve their bioavailability and CNS potential so that they can be explored for their utility in inflammatory CNS disorders (for a recent review on the topic see [51]).

MSK-1/2: Is this downstream target of p38 the brake on runaway inflammation?

Mitogen- and stress-activated kinases 1 and 2 (MSK1 and 2) are two related kinases that are activated downstream of p38 and Erk1/2 [52]. Both MSK1 and MSK2 are important for CREB and ATF1 phosphorylation in response to stress (UV or anisomycin) but not in response to mitogen (epidermal growth factor) [53]. MSK-1 can act on NF κ B to cause prolonged activation [39]. The single deletion of either MSK isoform caused only a small reduction in the cellular response to either stress-induced or mitogen-induced stimuli [53]. However, in the absence of both MSK1 and MSK2, there was a dramatic reduction in phosphorylation of CREB and ATF1. Two CREB-dependent immediate early genes c-fos and junB also showed a blunted induction in response to anisomycin (a strong p38 activator) in the MSK1/2 KO fibroblasts, but not in the single MSK deletion [53].

As a downstream target of p38, MSK1/2 was postulated to be important for the production of proinflammatory cytokines, reminiscent of the role of MK2. This question was addressed directly by Ananieva et al. [20, 54], where they found that MSK1/2 has anti-inflammatory activities and helps restrain inflammation. Using the MSK1/2 double KO mice, they found that there was a dramatic 20 to 60-fold increase in TNF α , IL-6 and IL-12 in bone marrow derived macrophages stimulated with LPS. Compared

to the WT bone marrow derived macrophages, there was also a decrease in the anti-inflammatory cytokine IL-10 in the MSK1/2 double KO macrophages [20, 54]. Using cells obtained from a triple KO mouse for MSK1/2 and IL-10, Ananieva et al. [20, 54] showed that the increase in TNF and IL-12 was directly dependent on MSK1/2. The triple KO mouse macrophages had greater IL-12 and TNF than macrophages from the single IL-10 KO. However, the increase in IL-6 was dependent on IL-10, as both the triple KO and the single IL-10 KO had similar levels of IL-6. When tested *in vivo*, a similar increase in TNF, IL-6, IL-12 and a decrease in IL-10 were found in the serum of mice injected with LPS [54]. In addition to IL-10, MSK1/2 is also important for the production of IL-1 receptor antagonist (IL-1ra) in response to LPS, partially independent of IL-10 [55]. Finally, MSK1/2 was also shown to increase DUSP1 transcription in macrophages, which can promote p38 dephosphorylation and deactivation [20, 54].

While the role of MSK1/2 has not been tested in any CNS-relevant *in vivo* models, it has been shown that MSK1 is expressed in the CNS. Webber et al. [56] measured MSK1 phosphorylation in the hippocampus and cortex of 10 AD and 9 non-AD patients, and found that phospho-MSK1 was dramatically increased in the cytoplasm of the pyramidal neurons in the AD brain. This was true for MSK-1 phosphorylated at Thr581 and MSK1 phosphorylated at Ser376, although the staining patterns for the two phospho-specific antibodies did not overlap. In the non-AD samples, there appeared to be no phospho-MSK-1 staining. Interestingly, the pattern of MSK-1 staining co-localized with phospho-ERK and not with phospho-p38. One caveat is that the staining was done on adjacent sections and not using confocal microscopy, so it is difficult to determine how much colocalization occurred in these samples [56].

By gene deletion studies, the p38 substrate MSK1/2 has clearly been shown to have anti-inflammatory properties (Fig.1). However, almost two decades of research have shown that p38 inhibitors are able to suppress the production of pro-inflammatory cytokines. One possible explanation for these seemingly contradictory findings could come from the way MSK1/2 suppresses inflammation, which can be by acting on p38 (Fig.2). Therefore if p38 is being blocked with a compound then suppression of MSK1/2 has a negated impact. The exception would be the decrease in IL-10, which suppresses

inflammation via a Jak/Stat mediated mechanism (for a recent review see [57]). Pharmacological inhibition typically does not completely inhibit the biological target the way a gene deletion does, which must be considered in interpreting data obtained from these two kinds of studies. In addition, in considering p38-driven signaling pathways as potential therapeutic targets, these studies provide another example of the importance of the therapeutic window. It will be important in the future to determine the appropriate window of responsiveness where there is adequate suppression of the pro-inflammatory responses while maintaining some of the anti-inflammatory responsiveness as well.

Is p38 MAPK pathway a potential therapeutic target for CNS inflammatory disorders?

The p38 α MAPK is one of the most well established therapeutic targets for peripheral inflammatory diseases, and several small molecule p38 α inhibitors are currently in human clinical trials for diseases such as rheumatoid arthritis, asthma, atherosclerosis, and acute lung injury [58]. There has also been a move in recent years to using other members of the MAPK pathway as therapeutic targets, such as MK2 or DUSPs (for recent review see [59, 60]). As we discuss in this review, there is extensive evidence that the p38 MAPK signaling cascade contributes to the cytokine overproduction in disease. Cytokine overproduction contributes to the neurodegenerative sequelae seen in a variety of neurodegenerative disorders. In considering whether a protein of interest should be explored as a potential therapeutic target for CNS inflammatory disorders, two important questions must be addressed. First, is the target present in the right time and place; i.e., is there evidence that it is involved in the disease process? Second, and more importantly, is the target a “druggable target”? In other words, can the target be modulated with a small molecule compound and achieve a therapeutically relevant outcome without untoward adverse effects? Examination of the available literature suggests that p38 α fulfills these two criteria for several CNS disorders.

Highlights from the AD field are presented that exemplify the evidence. To the first question – **right time and place**? For p38 α there is extensive evidence to address this question in AD. **Right time:** Stimulation of glial cell cultures with A β 1-42 induces p38 MAPK activation and subsequent induction of

proinflammatory cytokines [61-66]. Activation of neuronal p38 MAPK by RAGE contributes to translocation of A β into neurons, A β -induced impairment of cortical LTP, deficits in kinesin-dependent fast axonal transport, and A β -induced neurotoxicity [67-70]. The p38 MAPK can phosphorylate tau, *in vitro*, at AD-relevant amino acid residues [71-75], and is involved in cytokine-induced changes in tau phosphorylation and synaptic damage [76]. **Right place:** The p38 MAPK pathway is activated in the early stages of patients with AD and localizes to both neurons and glia [72,73,77-81]. Increases in activated p38 MAPK are seen coincident with AD-relevant pathology endpoints in animal models, such as increases in amyloid deposition, tau phosphorylation and loss of synaptophysin [64-66]. To the second question – *is it a druggable target?* A selective, orally active, brain-penetrant p38 α inhibitor

restores excessive glia proinflammatory cytokine production back towards normal and attenuates neurologic dysfunction in an AD-relevant mouse model, without observable adverse effects [26]. Altogether, p38 α MAPK is emerging as an attractive target for CNS inflammatory disorders. Current drug discovery efforts are building on this mechanistic knowledge in attempts to develop potential neurodegenerative disease-modifying therapeutics that target this critical gene-regulating protein kinase. Less evidence is currently available for MK2. As a direct downstream target of p38 α , MK2 would appear to be a logical suspect to investigate. It is still open to debate if MK2 is a good therapeutic target for CNS inflammatory disorders, but the odds are in favor of MK2 becoming an attractive avenue of research in the future.

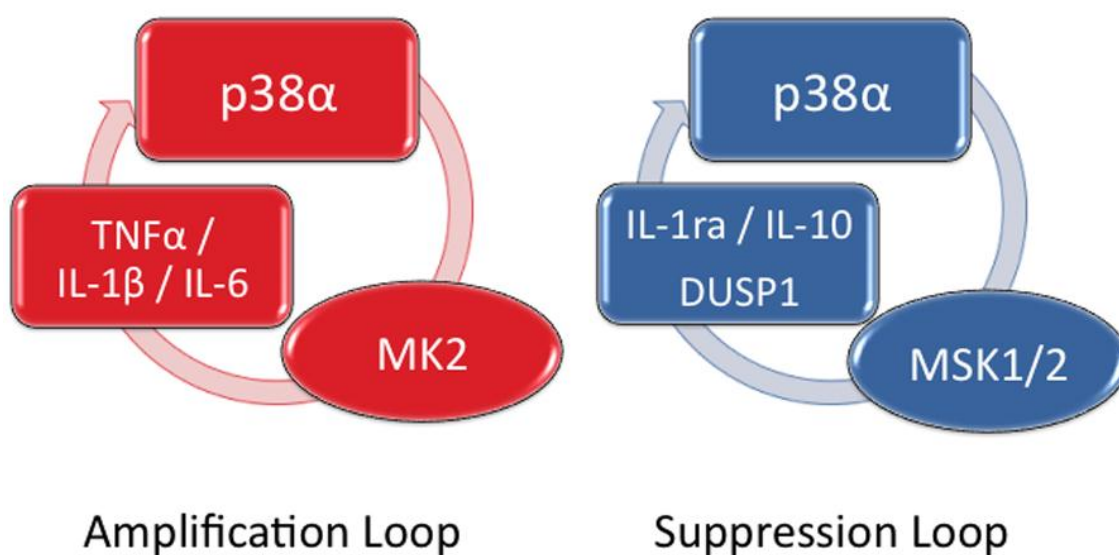


Figure 2: Feedback mechanisms regulating p38 α . Inflammatory signals, such as TNF α and IL-1 β , can activate the p38 pathway. Two downstream kinases of p38 are MK2 and MSK1/2. MK2 can promote an increase in TNF α , IL-1 β and IL-6. These pro-inflammatory cytokines can then feed back onto the cell, activating the same pathway and amplifying the inflammation. MSK1/2 produces the phosphatase DUSP1, which dephosphorylates and deactivates p38. Two antiinflammatory cytokines IL-1ra and IL-10 are also regulated by MSK1/2 activation. These cytokines have the potential to antagonize the effects of IL-1 β and suppress the inflammatory response.

Conclusion

Activation of the innate immune system occurs following pathogen invasion, chemical irritation, and mechanical trauma. The acute inflammatory

response limits ongoing infection and compartmentalizes damaged tissue. Failure to limit excessive activation of the inflammatory response produces acute disease such as septic shock, which can be fatal. Failure to temporally limit the

inflammatory response produces chronic disease, such as arthritis, asthma, atherosclerosis and age-related neurodegenerative diseases. Many kinases are involved in the inflammatory response. This review has focused on the role of the p38 MAPK family of kinases and the downstream kinases MK2 and MSK1/2 in cytokine production. Figure 1 shows a very simplified flow diagram of p38 α signaling, with figure 2 showing how p38 α through MK2 and MSK1/2 can affect cytokine production.

Although there has been extensive effort to understand and exploit the p38 MAPK pathway as therapeutic targets for peripheral inflammatory disorders, the field is just beginning to understand how to effectively use our cumulative knowledge to develop strategies to intervene in CNS inflammatory disorders. Selective, CNS-penetrant, small molecule inhibitors of p38 α , which have the potential to be used to treat disorders of the CNS, are being developed. One example of a promising, selective p38 α inhibitor that has the ability to cross the blood-brain barrier is MW01-2069A-SRM [26]. Small molecule inhibitors of MK2 are still in the early phases of development, and their potential use in CNS disorders needs to be explored. The pivotal role of p38 MAPK pathways in the regulation of responses that lead to overproduction of CNS proinflammatory cytokines emphasizes the importance of these drug discovery efforts.

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